

PCT09

RAW SEQUENCE LISTING

DATE: 11/14/2001

PATENT APPLICATION: US/09/763,011A

TIME: 13:52:05

Input Set : A:\JAB-1415.txt

Output Set: N:\CRF3\11142001\I763011A.raw

**Does Not Comply
Corrected Diskette Needed**

5 <110> APPLICANT: Contreras, Roland
6 Nelissen, Bart
7 DeBacker, Marianne
8 Luyten, Walter
9 Viaene, Jasmine
10 Logghe, Marc George
13 <120> TITLE OF INVENTION: Drug Targets In Candida Albicans
16 <130> FILE REFERENCE: 50899/002
C--> 19 <140> CURRENT APPLICATION NUMBER: US/09/763,011A 0K
C--> 21 <141> CURRENT FILING DATE: 2001-02-14
24 <150> PRIOR APPLICATION NUMBER: 98310694.9
26 <151> PRIOR FILING DATE: 1998-12-23
29 <150> PRIOR APPLICATION NUMBER: 9817796.7
31 <151> PRIOR FILING DATE: 1998-08-14
34 <160> NUMBER OF SEQ ID NOS: 114
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238 20 25 30
242 Ile Ser Ile Leu Thr Asn His Phe Gln Ile Leu Lys Asp Leu Leu Pro
244 35 40 45
248 Tyr Ser Lys Thr Ser Lys Pro Gln Ile Lys Glu Ser Arg Pro Leu Ile
250 50 55 60
254 Lys Val Ser Arg Asp Gly Val Pro Ile Asn Phe His Arg Ala Pro Ala
256 65 70 75 80
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262 85 90 95
266 Lys Thr Met Val Leu Thr Glu Ile Lys Thr Ile Thr Glu Phe Ala Thr
268 100 105 110
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310 210 215 220
314 Val Pro Lys Glu Glu Val Met Val Ser His Asp Thr Ala Pro Ile Val
316 225 230 235 240
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322 245 250 255
326 Ser Lys His Lys Met Asn Gly Lys Asn Ile Leu Asn Asn Asn Pro Pro
328 260 265 270
332 Pro Asn Ser Val Leu Ile Val Pro Gln Phe Gln Phe His Glu Arg Met
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344 Asn Phe Pro Thr Thr Thr Ala Thr Ser Met Phe Asp Asn Phe Ile Leu
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350 Lys Asn Leu Val Asp Glu Thr Asp Ser Glu Ile Asp Ser Gly Glu Thr
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370           370           375           380
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388           420           425           430
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436 tatcaagaga atgggcaaaa gaacagtcaa ccgatcccta acacaaacac agagcataat 240
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484 tttaaggata tgtatgagag tttgcctgac cattcaagta aggcaacacc taaagacgat 1680
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490 agtatgttgg acatgcatgc gtcttctaaa attcaaagtc ttttacctcc acaaccgcca 1860
492 caaatgtcaa ttgatccttc tgtttccaag caagtgtggg ctaaatacgt tgatgcaatc 1920
494 ttgacttatc aaagagaatt tttcaattat aaaaaagtga ttgttcaata ccaaatggaa 1980
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498 ttggatactt tcaagcattg tttagaacaa gattatttgg ttatgagtga gtttaatgaa 2100
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526 20 25 30
530 Tyr His Pro Asp Lys Thr Pro Arg Arg Glu Asp His Glu Lys Phe Lys
532 35 40 45
536 Glu Ile Asn Ile Ala Tyr Glu Thr Ile Arg Asp Tyr Tyr Gln Glu Asn
538 50 55 60
542 Gly Gln Lys Asn Ser Gln Pro Ile Pro Asn Thr Asn Thr Glu His Asn
544 65 70 75 80
548 Ser His Gln Lys Pro His Tyr Asn Thr Gly Pro Tyr Ser Thr Tyr Arg
550 85 90 95
554 Phe Thr Thr Ser Ser Thr Thr Thr Asp Asn Thr Asn His Thr Gly His
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562 115 120 125
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572 Lys Lys Glu Leu Phe Gln Arg Gln Gln Ala Glu Glu Ala Gln Arg Lys
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580 165 170 175
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586 180 185 190
590 Gln Arg Gln Lys Glu His Glu Ala Lys Arg Asp Ile Arg Ile Gln Gln
592 195 200 205
596 Leu Ser Glu Gln Asp Ser Arg Ser Asn Gln Thr Lys Glu Glu Glu Glu
598 210 215 220
602 Val Phe Lys Lys Ala Arg Ser Thr Asn Ser Gly Ala Asp Glu Thr Gly
604 225 230 235 240
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09/763011A

Page 5 of 7B

<210> 13

<211> 302

<212> PRT

<213> Candida albicans

<220>

<221> Xaa = any amino acid at positions 58 and 185

**Does Not Comply
Corrected Diskette Needed**

1. Unknowns must be enumerated in fields 221 222 and 223.
2. Field 221 as "variant"
3. Field 222 as location "58 and 185"
4. Field 223 as particular amino proteins

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

DATE: 11/14/2001

PATENT APPLICATION: US/09/763,011A

TIME: 13:52:06

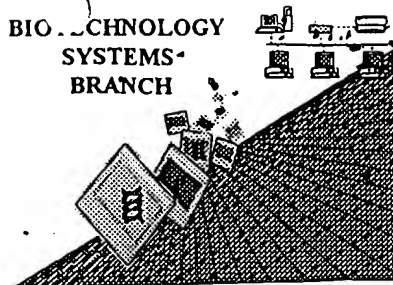
Input Set : A:\JAB-1415.txt

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L:21 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:1041 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:13
L:1067 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:13
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L:1067 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
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RAW SEQUENCE LISTING ERROR REPORT

BIO-TECHNOLOGY
SYSTEMS-
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09 763 011A

Source: PCT 09

Date Processed by STIC: 11/14/01

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>